

SEQUENCE LISTING

<110> Knutzon, Debbie

<120> POLYUNSATURATED FATTY ACIDS IN PLANTS

<130> MOCO.156.00US

<140> 09/330,235

<141> 1999-06-10

<150> 60/089,043

<151> 1998-06-12

<160> 22

<170> PatentIn version 3.0

<210> 1

<211> 1391

<212> DNA

<213> Caenorhabditis elegans

<400> 1

```

caagtttgag gtatgggtcgc tcattcctca gaaggggttat ccgccacggc tccggtcacc      60
ggcggagatg ttctggttga tgctcgtgca tctcttgaag aaaaggaggc tccacgtgat      120
gtgaatgcaa aactaaaca ggccaccact gaagagccac gcatccaatt accaactgtg      180
gatgctttcc gtcgtgcaat tccagcacac tgtttcgaaa gagatctcgt taaatcaatc      240
agatatttgg tgcaagactt tgcggcactc acaattctct actttgctct tccagctttt      300
gagtactttg gattgtttgg ttacttgggt tggaacattt ttatgggagt ttttggattc      360
gcgttggtcg tcgttggaca cgattgtctt catggatcat tctctgataa tcagaatctc      420
aatgatttca ttggacatat cgccttctca ccactcttct ctccatactt cccatggcag      480
aaaagtcaca agcttcacca tgctttcacc aaccacattg acaaagatca tggacacgtg      540
tggattcagg ataaggattg ggaagcaatg ccatcatgga aaagatggtt caatccaatt      600
ccattctctg gatggcttaa atgggtccca gtgtacactt tattcgggtt ctgtgatgga      660
tctcacttct ggccatactc ttcacttttt gttcgtaact ctgaccgtgt tcaatgtgta      720
atctctggaa tctgttgctg tgtgtgtgca tatattgctc taacaattgc tggatcatat      780
tccaattggt tctggtacta ttgggttcca ctttctttct tcggattgat gctcgtcatt      840
gttacctatt tgcaacatgt cgatgatgtc gctgaggtgt acgaggctga tgaatggagc      900
ttcgtccgtg gacaaacca aaccatcgat cgttactatg gactcggatt ggacacaacg      960
atgcaccata tcacagacgg acacgttgcc catcacttct tcaacaaaat cccacattac     1020

```

catctcatcg aagcaaccga aggtgtcaaa aaggtcttgg agccgttgtc cgacacccaa 1080
 tacgggtaca aatctcaagt gaactacgat ttctttgccc gtttcctgtg gttcaactac 1140
 aagctcgact atctcgttca caagaccgcc ggaatcatgc aattccgaac aactctcgag 1200
 gagaaggcaa aggccaagta aaagaatatc ccgtgccgtt ctagagtaca acaacaactt 1260
 ctgcgttttc accggttttg ctctaattgc aatttttctt tgttctatat atattttttt 1320
 gctttttaat tttattctct ctaaaaaact tctacttttc agtgcgttga atgcataaag 1380
 ccataactct t 1391

<210> 2
 <211> 402
 <212> PRT
 <213> Caenorhabditis elegans

<400> 2

Met Val Ala His Ser Ser Glu Gly Leu Ser Ala Thr Ala Pro Val Thr
 1 5 10 15
 Gly Gly Asp Val Leu Val Asp Ala Arg Ala Ser Leu Glu Glu Lys Glu
 20 25 30
 Ala Pro Arg Asp Val Asn Ala Asn Thr Lys Gln Ala Thr Thr Glu Glu
 35 40 45
 Pro Arg Ile Gln Leu Pro Thr Val Asp Ala Phe Arg Arg Ala Ile Pro
 50 55 60
 Ala His Cys Phe Glu Arg Asp Leu Val Lys Ser Ile Arg Tyr Leu Val
 65 70 75 80
 Gln Asp Phe Ala Ala Leu Thr Ile Leu Tyr Phe Ala Leu Pro Ala Phe
 85 90 95
 Glu Tyr Phe Gly Leu Phe Gly Tyr Leu Val Trp Asn Ile Phe Met Gly
 100 105 110
 Val Phe Gly Phe Ala Leu Phe Val Val Gly His Asp Cys Leu His Gly
 115 120 125
 Ser Phe Ser Asp Asn Gln Asn Leu Asn Asp Phe Ile Gly His Ile Ala
 130 135 140
 Phe Ser Pro Leu Phe Ser Pro Tyr Phe Pro Trp Gln Lys Ser His Lys
 145 150 155 160
 Leu His His Ala Phe Thr Asn His Ile Asp Lys Asp His Gly His Val
 165 170 175
 Trp Ile Gln Asp Lys Asp Trp Glu Ala Met Pro Ser Trp Lys Arg Trp

<400> 4
caucaucauc augcggccgc ttacttggcc ttgcctt 38

<210> 5
<211> 32
<212> DNA
<213> synthetic polylinker

<400> 5
tcgacctgca ggaagcttgc ggccgcggat cc 32

<210> 6
<211> 32
<212> DNA
<213> synthetic polylinker

<400> 6
tcgaggatcc gcggccgcaa gcttcctgca gg 32

<210> 7
<211> 1353
<212> DNA
<213> Brassica napus

<400> 7
aatccatcaa acctttattc accacatttc actgaaaggc cacacatcta gagagagaaa 60
cttcgtccaa atctctctct ccagcgatgg ttgttgctat ggaccagcgc agcaatgtta 120
acggagattc cggtgcccgg aaggaagaag ggtttgatcc aagcgcacaa ccaccgttta 180
agatcggaga tataagggcg gcgattccta agcattgctg ggtgaagagt cctttgagat 240
ctatgagcta cgtcaccaga gacattttcg ccgtcgcggc tctggccatg gccgccgtgt 300
attttgatag ctggttcctc tggccactct actgggttgc ccaaggaacc cttttctggg 360
ccatcttcgt tcttggccac gactgtggac atgggagttt ctacagacatt cctctgctga 420
acagtgtggt tggtcacatt cttcattcat tcatcctcgt tccttaccat ggttggagaa 480
taagccatcg gacacaccac cagaaccatg gccatgttga aaacgacgag tcttgggttc 540
cgttgccaga aaagttgtac aagaacttgc cccatagtac tcggatgctc agatacactg 600
tcctctgcc catgctcgtt taccgatct atctgtggtg cagaagtcct ggaaaagaag 660
ggtcacattt taaccatac agtagtttat ttgctccaag cgagaggaag cttattgcaa 720
cttcaactac ttgctggtec ataatgttgg ccactcttgt ttatctatcg ttctcgttg 780
atccagtcac agttctcaaa gtctatggcg ttccttacat tatctttgtg atgtgggttg 840

acgctgtcac gtacttgcac catcatggtc acgatgagaa gttgccttgg tacagaggca 900
 aggaatggag ttattttacgt ggaggattaa caactattga tagagattac ggaatcttca 960
 acaacatcca tcacgacatt ggaactcacg tgatccatca tcttttccca caaatccctc 1020
 actatcactt ggtcgaagcc acgagagcag ctaaacaatgt gttaggaaga tactacagag 1080
 agccgaagac gtcaggagca ataccgattc acttggtgga gagtttggtc gcaagtatta 1140
 aaaaagatca ttacgtcagt gacactgggtg atattgtctt ctacgagaca gatccagatc 1200
 tctacgttta tgcttctgac aaatctaaaa tcaattaact tttcttccca gctctattag 1260
 gaataaacac tccttctctt ttacttattt gtttctgctt taagtttaaa atgtactcgt 1320
 gaaacctttt ttttattaat gtattttacgt tac 1353

<210> 8
 <211> 383
 <212> PRT
 <213> Brassica napus

<400> 8

Met Val Val Ala Met Asp Gln Arg Ser Asn Val Asn Gly Asp Ser Gly
 1 5 10 15
 Ala Arg Lys Glu Glu Gly Phe Asp Pro Ser Ala Gln Pro Pro Phe Lys
 20 25 30
 Ile Gly Asp Ile Arg Ala Ala Ile Pro Lys His Cys Trp Val Lys Ser
 35 40 45
 Pro Leu Arg Ser Met Ser Tyr Val Thr Arg Asp Ile Phe Ala Val Ala
 50 55 60
 Ala Leu Ala Met Ala Ala Val Tyr Phe Asp Ser Trp Phe Leu Trp Pro
 65 70 75 80
 Leu Tyr Trp Val Ala Gln Gly Thr Leu Phe Trp Ala Ile Phe Val Leu
 85 90 95
 Gly His Asp Cys Gly His Gly Ser Phe Ser Asp Ile Pro Leu Leu Asn
 100 105 110
 Ser Val Val Gly His Ile Leu His Ser Phe Ile Leu Val Pro Tyr His
 115 120 125
 Gly Trp Arg Ile Ser His Arg Thr His His Gln Asn His Gly His Val
 130 135 140
 Glu Asn Asp Glu Ser Trp Val Pro Leu Pro Glu Lys Leu Tyr Lys Asn
 145 150 155 160
 Leu Pro His Ser Thr Arg Met Leu Arg Tyr Thr Val Pro Leu Pro Met

165										170					175				
Leu	Ala	Tyr	Pro	Ile	Tyr	Leu	Trp	Tyr	Arg	Ser	Pro	Gly	Lys	Glu	Gly				
			180					185					190						
Ser	His	Phe	Asn	Pro	Tyr	Ser	Ser	Leu	Phe	Ala	Pro	Ser	Glu	Arg	Lys				
		195					200					205							
Leu	Ile	Ala	Thr	Ser	Thr	Thr	Cys	Trp	Ser	Ile	Met	Leu	Ala	Thr	Leu				
	210					215					220								
Val	Tyr	Leu	Ser	Phe	Leu	Val	Asp	Pro	Val	Thr	Val	Leu	Lys	Val	Tyr				
225					230					235					240				
Gly	Val	Pro	Tyr	Ile	Ile	Phe	Val	Met	Trp	Leu	Asp	Ala	Val	Thr	Tyr				
				245					250					255					
Leu	His	His	His	Gly	His	Asp	Glu	Lys	Leu	Pro	Trp	Tyr	Arg	Gly	Lys				
			260					265					270						
Glu	Trp	Ser	Tyr	Leu	Arg	Gly	Gly	Leu	Thr	Thr	Ile	Asp	Arg	Asp	Tyr				
		275					280					285							
Gly	Ile	Phe	Asn	Asn	Ile	His	His	Asp	Ile	Gly	Thr	His	Val	Ile	His				
	290					295					300								
His	Leu	Phe	Pro	Gln	Ile	Pro	His	Tyr	His	Leu	Val	Asp	Ala	Thr	Arg				
305					310					315					320				
Ala	Ala	Lys	His	Val	Leu	Gly	Arg	Tyr	Tyr	Arg	Glu	Pro	Lys	Thr	Ser				
				325					330					335					
Gly	Ala	Ile	Pro	Ile	His	Leu	Val	Glu	Ser	Leu	Val	Ala	Ser	Ile	Lys				
			340					345					350						
Lys	Asp	His	Tyr	Val	Ser	Asp	Thr	Gly	Asp	Ile	Val	Phe	Tyr	Glu	Thr				
		355					360					365							
Asp	Pro	Asp	Leu	Tyr	Val	Tyr	Ala	Ser	Asp	Lys	Ser	Lys	Ile	Asn					
	370					375					380								

<210> 9
 <211> 40
 <212> DNA
 <213> synthetic primer

<400> 9
 cuacuacuac uagagctcag cgatggttgt tgctatggac

40

<210> 10
 <211> 37
 <212> DNA
 <213> synthetic primer

<400> 10
 caucaucauc augaattctt aattgatttt agatttg

37

<210> 11
 <211> 1482
 <212> DNA
 <213> Mortierella alpina

<400> 11
 gcttcctcca gttcatcctc catttcgcca cctgcattct ttacgaccgt taagcaagat 60
 gggaacggac caaggaaaaa ctttcacctg ggaagagctg gcggcccata acaccaagga 120
 cgacctactc ttggccatcc gcggcagggg gtacgatgtc acaaagttct tgagccgcca 180
 tcctgggtgga gtggacactc tctgtctcgg agctggccga gatgttactc cggctcttga 240
 gatgtatcac gcgtttgggg ctgcagatgc cattatgaag aagtactatg tcggtacact 300
 ggtctcgaat gagctgcccc tcttccccga gccaacgggtg ttccacaaaa ccatcaagac 360
 gagagtcgag ggctacttta cggatcggaa cattgatccc aagaatagac cagagatctg 420
 gggacgatac gctcttatct ttggatcctt gatcgcttcc tactacgcgc agctctttgt 480
 gcctttcggt gtcgaaacgca catggcttca ggtgggtgtt gcaatcatca tgggatttgc 540
 gtgcgcacaa gtccgactca accctcttca tgatgcgtct cacttttcag tgaccacaaa 600
 ccccaactgtc tggaagattc tgggagccac gcacgacttt ttcaacggag catcgtaact 660
 ggtgtggatg taccaacata tgctcggcca tcaccctac accaacattg ctggagcaga 720
 tcccgaactg tcgacgtctg agcccgatgt tcgtcgtatc aagcccaacc aaaagtgggt 780
 tgtcaaccac atcaaccagc acatgtttgt tcttttctg tacggactgc tggcgttcaa 840
 ggtgcgcatt caggacatca acattttgta ctttgtcaag accaatgacg ctattcgtgt 900
 caatcccatc tcgacatggc aactgtgat gttctggggc ggcaaggctt tctttgtctg 960
 gtatcgctg attgttcccc tgcagtatct gccctgggc aagggtgctgc tcttgttcac 1020
 ggtcgcggac atgggtgtcgt cttactgggt ggcgtgacc ttccaggcga accacgttgt 1080
 tgaggaagtt cagtggcgt tgctgacga gaacgggatc atccaaaagg actgggcagc 1140
 tatgcaggtc gagactacgc aggattacgc acacgattcg cacctctgga ccagcatcac 1200
 tggcagcttg aactaccagg ctgtgcacca tctgttcccc aacgtgtcgc agcaccatta 1260
 tcccgatatt ctggccatca tcaagaacac ctgcagcgag tacaaggttc cataccttgt 1320
 caaggatacg ttttggcaag catttgcttc acatttggag cacttgctg tctttggact 1380
 ccgtcccaag gaagagtaga agaaaaaag cgccgaatga agtattgccc cctttttctc 1440
 caagaatggc aaaaggagat caagtggaca ttctctatga ag 1482

<210> 12
 <211> 446
 <212> PRT
 <213> Mortierella alpina

<400> 12

Met Gly Thr Asp Gln Gly Lys Thr Phe Thr Trp Glu Glu Leu Ala Ala
 1 5 10 15
 His Asn Thr Lys Asp Asp Leu Leu Leu Ala Ile Arg Gly Arg Val Tyr
 20 25 30
 Asp Val Thr Lys Phe Leu Ser Arg His Pro Gly Gly Val Asp Thr Leu
 35 40 45
 Leu Leu Gly Ala Gly Arg Asp Val Thr Pro Val Phe Glu Met Tyr His
 50 55 60
 Ala Phe Gly Ala Ala Asp Ala Ile Met Lys Lys Tyr Tyr Val Gly Thr
 65 70 75 80
 Leu Val Ser Asn Glu Leu Pro Ile Phe Pro Glu Pro Thr Val Phe His
 85 90 95
 Lys Thr Ile Lys Thr Arg Val Glu Gly Tyr Phe Thr Asp Arg Asn Ile
 100 105 110
 Asp Pro Lys Asn Arg Pro Glu Ile Trp Gly Arg Tyr Ala Leu Ile Phe
 115 120 125
 Gly Ser Leu Ile Ala Ser Tyr Tyr Ala Gln Leu Phe Val Pro Phe Val
 130 135 140
 Val Glu Arg Thr Trp Leu Gln Val Val Phe Ala Ile Ile Met Gly Phe
 145 150 155 160
 Ala Cys Ala Gln Val Gly Leu Asn Pro Leu His Asp Ala Ser His Phe
 165 170 175
 Ser Val Thr His Asn Pro Thr Val Trp Lys Ile Leu Gly Ala Thr His
 180 185 190
 Asp Phe Phe Asn Gly Ala Ser Tyr Leu Val Trp Met Tyr Gln His Met
 195 200 205
 Leu Gly His His Pro Tyr Thr Asn Ile Ala Gly Ala Asp Pro Asp Val
 210 215 220
 Ser Thr Ser Glu Pro Asp Val Arg Arg Ile Lys Pro Asn Gln Lys Trp
 225 230 235 240
 Phe Val Asn His Ile Asn Gln His Met Phe Val Pro Phe Leu Tyr Gly
 245 250 255

Leu Leu Ala Phe Lys Val Arg Ile Gln Asp Ile Asn Ile Leu Tyr Phe
 260 265 270

Val Lys Thr Asn Asp Ala Ile Arg Val Asn Pro Ile Ser Thr Trp His
 275 280 285

Thr Val Met Phe Trp Gly Gly Lys Ala Phe Phe Val Trp Tyr Arg Leu
 290 295 300

Ile Val Pro Leu Gln Tyr Leu Pro Leu Gly Lys Val Leu Leu Leu Phe
 305 310 315 320

Thr Val Ala Asp Met Val Ser Ser Tyr Trp Leu Ala Leu Thr Phe Gln
 325 330 335

Ala Asn His Val Val Glu Glu Val Gln Trp Pro Leu Pro Asp Glu Asn
 340 345 350

Gly Ile Ile Gln Lys Asp Trp Ala Ala Met Gln Val Glu Thr Thr Gln
 355 360 365

Asp Tyr Ala His Asp Ser His Leu Trp Thr Ser Ile Thr Gly Ser Leu
 370 375 380

Asn Tyr Gln Ala Val His His Leu Phe Pro Asn Val Ser Gln His His
 385 390 395 400

Tyr Pro Asp Ile Leu Ala Ile Ile Lys Asn Thr Cys Ser Glu Tyr Lys
 405 410 415

Val Pro Tyr Leu Val Lys Asp Thr Phe Trp Gln Ala Phe Ala Ser His
 420 425 430

Leu Glu His Leu Arg Val Leu Gly Leu Arg Pro Lys Glu Glu
 435 440 445

<210> 13
 <211> 39
 <212> DNA
 <213> synthetic primer

<400> 13
 cuacuacuac uactcgagca agatgggaac ggaccaagg

39

<210> 14
 <211> 39
 <212> DNA
 <213> synthetic primer

<400> 14
 caucauac auctcgagct actcttcctt gggacggag

39

<210> 15
 <211> 47
 <212> DNA

<213> synthetic primer

<400> 15

cuacuacuac uatctagact cgagaccatg gctgctgctc cagtgtg

47

<210> 16

<211> 40

<212> DNA

<213> synthetic primer

<400> 16

caucaucauc auaggcctcg agttactgcg ccttaccat

40

<210> 17

<211> 1617

<212> DNA

<213> Mortierella alpina

<400> 17

cgacactcct tccttcttct caccgcctct agtccccttc aacccccctc tttgacaaag 60
acaacaaacc atggctgctg ctcccagtggt gaggacgttt actcggggccg aggtttttgaa 120
tgccgagggt ctgaatgagg gcaagaagga tgccgaggca cccttcttga tgatcatcga 180
caacaagggt tacgatgtcc gcgagttcgt ccctgatcat cccggtggaa gtgtgattct 240
cacgcacgtt ggcaaggacg gcactgacgt ctttgacact tttcaccctg aggtgcttg 300
ggagactcct gccaaactttt acgttggtga tattgacgag agcgaccgcg atatcaagaa 360
tgatgacttt gcggccgagg tccgcaagct gcgtacctg ttccagtctc ttggttacta 420
cgattcttcc aaggcatact acgccttcaa ggtctcgttc aacctctgca tctggggttt 480
gtcgacggtc attgtggcca agtggggcca gacctcgacc ctgcgcaacg tgctctcggc 540
tgcgcttttg ggtctgttct ggcagcagtg cggatggttg gctcacgact ttttgcac 600
ccaggtcttc caggaccgtt tctgggggtga tcttttcggc gccttcttgg gaggtgtctg 660
ccagggtctc tegtctctgt ggtggaagga caagcacaac actcaccacg ccgcccccaa 720
cgtccacggc gaggatcccg acattgacac ccacctctg ttgacctgga gtgagcatgc 780
gttgagatg ttctcggatg tcccagatga ggagctgacc cgcagtgtgt cgcgtttcat 840
ggtcctgaac cagacctggt tttacttccc cattctctcg tttgcccgtc tctcctggtg 900
cctccagtcc attctctttg tgctgcctaa cggtcaggcc cacaagccct cgggcgcgcg 960
tgtgcccac tcgttggtcg agcagctgtc gcttgcgatg cactggacct ggtacctcgc 1020
caccatgttc ctgttcatca aggatcccg caacatgctg gtgtactttt tgggtgtcgca 1080

ggcggtgtgc ggaaacttgt tggcgatcgt gttctcgtc aaccacaacg gtatgcctgt 1140
 gatctcgaag gaggaggcgg tcgatatgga tttcttcacg aagcagatca tcacgggtcg 1200
 tgatgtccac ccgggtctat ttgccaaactg gttcacgggt ggattgaact atcagatcga 1260
 gcaccacttg ttccttcga tgctcgcca caacttttca aagatccagc ctgctgtcga 1320
 gaccctgtgc aaaaagtaca atgtccgata ccacaccacc ggtatgatcg agggaaactgc 1380
 agaggtcttt agccgtctga acgaggtctc caaggctgcc tccaagatgg gtaaggcgca 1440
 gtaaaaaaaaa aaacaaggac gttttttttc gccagtgcct gtgctgtgc ctgcttccct 1500
 tgtcaagtcg agcgtttctg gaaaggatcg ttcagtgcag tatcatcatt ctctttttac 1560
 cccccgtca tatctcatte atttctctta ttaaacaact tgttcccccc ttcaccg 1617

<210> 18
 <211> 457
 <212> PRT
 <213> Mortierella alpina

<400> 18

Met Ala Ala Ala Pro Ser Val Arg Thr Phe Thr Arg Ala Glu Val Leu
 1 5 10 15

Asn Ala Glu Ala Leu Asn Glu Gly Lys Lys Asp Ala Glu Ala Pro Phe
 20 25 30

Leu Met Ile Ile Asp Asn Lys Val Tyr Asp Val Arg Glu Phe Val Pro
 35 40 45

Asp His Pro Gly Gly Ser Val Ile Leu Thr His Val Gly Lys Asp Gly
 50 55 60

Thr Asp Val Phe Asp Thr Phe His Pro Glu Ala Ala Trp Glu Thr Leu
 65 70 75 80

Ala Asn Phe Tyr Val Gly Asp Ile Asp Glu Ser Asp Arg Asp Ile Lys
 85 90 95

Asn Asp Asp Phe Ala Ala Glu Val Arg Lys Leu Arg Thr Leu Phe Gln
 100 105 110

Ser Leu Gly Tyr Tyr Asp Ser Ser Lys Ala Tyr Tyr Ala Phe Lys Val
 115 120 125

Ser Phe Asn Leu Cys Ile Trp Gly Leu Ser Thr Val Ile Val Ala Lys
 130 135 140

Trp Gly Gln Thr Ser Thr Leu Ala Asn Val Leu Ser Ala Ala Leu Leu
 145 150 155 160

Gly Leu Phe Trp Gln Gln Cys Gly Trp Leu Ala His Asp Phe Leu His

165	170	175
His Gln Val Phe Gln Asp Arg Phe Trp Gly Asp Leu Phe Gly Ala Phe 180 185 190		
Leu Gly Gly Val Cys Gln Gly Phe Ser Ser Ser Trp Trp Lys Asp Lys 195 200 205		
His Asn Thr His His Ala Ala Pro Asn Val His Gly Glu Asp Pro Asp 210 215 220		
Ile Asp Thr His Pro Leu Leu Thr Trp Ser Glu His Ala Leu Glu Met 225 230 235 240		
Phe Ser Asp Val Pro Asp Glu Glu Leu Thr Arg Met Trp Ser Arg Phe 245 250 255		
Met Val Leu Asn Gln Thr Trp Phe Tyr Phe Pro Ile Leu Ser Phe Ala 260 265 270		
Arg Leu Ser Trp Cys Leu Gln Ser Ile Leu Phe Val Leu Pro Asn Gly 275 280 285		
Gln Ala His Lys Pro Ser Gly Ala Arg Val Pro Ile Ser Leu Val Glu 290 295 300		
Gln Leu Ser Leu Ala Met His Trp Thr Trp Tyr Leu Ala Thr Met Phe 305 310 315 320		
Leu Phe Ile Lys Asp Pro Val Asn Met Leu Val Tyr Phe Leu Val Ser 325 330 335		
Gln Ala Val Cys Gly Asn Leu Leu Ala Ile Val Phe Ser Leu Asn His 340 345 350		
Asn Gly Met Pro Val Ile Ser Lys Glu Glu Ala Val Asp Met Asp Phe 355 360 365		
Phe Thr Lys Gln Ile Ile Thr Gly Arg Asp Val His Pro Gly Leu Phe 370 375 380		
Ala Asn Trp Phe Thr Gly Gly Leu Asn Tyr Gln Ile Glu His His Leu 385 390 395 400		
Phe Pro Ser Met Pro Arg His Asn Phe Ser Lys Ile Gln Pro Ala Val 405 410 415		
Glu Thr Leu Cys Lys Lys Tyr Asn Val Arg Tyr His Thr Thr Gly Met 420 425 430		
Ile Glu Gly Thr Ala Glu Val Phe Ser Arg Leu Asn Glu Val Ser Lys 435 440 445		
Ala Ala Ser Lys Met Gly Lys Ala Gln 450 455		

<210> 19

<211> 1488
 <212> DNA
 <213> Mortierella alpina

<400> 19
 gtccccgtgc gctgtcggca caccocatcc tccctcgctc cctctcggtt tgtccttggc 60
 ccaccgtctc tccctccacc tccgagacga ctgcaactgt aatcaggaac cgacaaatac 120
 acgattttctt ttactcagc accaactcaa aatcctcaac cgcaaccctt tttcaggatg 180
 gcacctccca acactatcga tgccgggttg acccagcgtc atatcagcac ctccggcccca 240
 aactcggcca agcctgcctt cgagcgcaac taccagctcc ccgagttcac catcaaggag 300
 atccgagagt gcatccctgc ccactgcttt gagcgctccg gtctccgtgg tctctgccac 360
 gttgccatcg atctgacttg ggcgtcgctc ttgttccttg ctgcgacca gatcgacaag 420
 tttgagaatc ccttgatccg ctatttgccc tggcctgttt actggatcat gcagggtatt 480
 gtctgcaccg gtgtctgggt gctggctcac gagtgtggtc atcagtcctt ctgcacctcc 540
 aagaccctca acaacacagt tggttggatc ttgcactcga tgctcttggg cccctaccac 600
 tccctggagaa tctcgcactc gaagcaccac aaggccactg gccatatgac caaggaccag 660
 gtctttgtgc ccaagaccgc ctcccagggt ggcttgccct ccaaggagaa cgctgctgct 720
 gccgttcagg aggaggacat gtccgtgcac ctggatgagg aggcctccat tgtgactttg 780
 ttctggatgg tgatccagtt cttgttcgga tggcccgct acctgattat gaacgcctct 840
 ggccaagact acggccgctg gacctcgac ttccacacgt actcgcccat ctttgagccc 900
 cgcaactttt tcgacattat tatctcgac ctccgtgtgt tggctgcctt cggctgcctg 960
 atctatgcct ccattgcagtt gtcgtctctg accgtcacca agtactatat tgtccctac 1020
 ctctttgtca acttttgggt ggtcctgac accctcttgc agcacaccga tccaagctg 1080
 cccattacc gcgaggggtg ctggaatttc cagcgtggag ctctttgcac cgttgaccgc 1140
 tcgtttggca agttcttggg ccatatgttc caaggcattg tccacacca tgtggcccat 1200
 cacttggtct cgcaaagcc gttctaccat gctgaggaag ctacctatca tctcaagaaa 1260
 ctgctgggag agtactatgt gtacgacca tccccgatcg tcgttgcggt ctggaggctg 1320
 ttccgtgagt gccgattcgt ggaggatcag ggagacgtgg tctttttcaa gaagtaaaaa 1380
 aaaagacaat ggaccacaca caaccttgct tctacagacc tacgtatcat gtagccatac 1440
 cacttcataa aagaacatga gctctagagg cgtgtcatte gcgcctcc 1488

<210> 20

<211> 399
 <212> PRT
 <213> Mortierella alpina

<400> 20

Met	Ala	Pro	Pro	Asn	Thr	Ile	Asp	Ala	Gly	Leu	Thr	Gln	Arg	His	Ile	1	5	10	15
Ser	Thr	Ser	Ala	Pro	Asn	Ser	Ala	Lys	Pro	Ala	Phe	Glu	Arg	Asn	Tyr	20	25	30	
Gln	Leu	Pro	Glu	Phe	Thr	Ile	Lys	Glu	Ile	Arg	Glu	Cys	Ile	Pro	Ala	35	40	45	
His	Cys	Phe	Glu	Arg	Ser	Gly	Leu	Arg	Gly	Leu	Cys	His	Val	Ala	Ile	50	55	60	
Asp	Leu	Thr	Trp	Ala	Ser	Leu	Leu	Phe	Leu	Ala	Ala	Thr	Gln	Ile	Asp	65	70	75	80
Lys	Phe	Glu	Asn	Pro	Leu	Ile	Arg	Tyr	Leu	Ala	Trp	Pro	Val	Tyr	Trp	85	90	95	
Ile	Met	Gln	Gly	Ile	Val	Cys	Thr	Gly	Val	Trp	Val	Leu	Ala	His	Glu	100	105	110	
Cys	Gly	His	Gln	Ser	Phe	Ser	Thr	Ser	Lys	Thr	Leu	Asn	Asn	Thr	Val	115	120	125	
Gly	Trp	Ile	Leu	His	Ser	Met	Leu	Leu	Val	Pro	Tyr	His	Ser	Trp	Arg	130	135	140	
Ile	Ser	His	Ser	Lys	His	His	Lys	Ala	Thr	Gly	His	Met	Thr	Lys	Asp	145	150	155	160
Gln	Val	Phe	Val	Pro	Lys	Thr	Arg	Ser	Gln	Val	Gly	Leu	Pro	Pro	Lys	165	170	175	
Glu	Asn	Ala	Ala	Ala	Ala	Val	Gln	Glu	Glu	Asp	Met	Ser	Val	His	Leu	180	185	190	
Asp	Glu	Glu	Ala	Pro	Ile	Val	Thr	Leu	Phe	Trp	Met	Val	Ile	Gln	Phe	195	200	205	
Leu	Phe	Gly	Trp	Pro	Ala	Tyr	Leu	Ile	Met	Asn	Ala	Ser	Gly	Gln	Asp	210	215	220	
Tyr	Gly	Arg	Trp	Thr	Ser	His	Phe	His	Thr	Tyr	Ser	Pro	Ile	Phe	Glu	225	230	235	240
Pro	Arg	Asn	Phe	Phe	Asp	Ile	Ile	Ile	Ser	Asp	Leu	Gly	Val	Leu	Ala	245	250	255	
Ala	Leu	Gly	Ala	Leu	Ile	Tyr	Ala	Ser	Met	Gln	Leu	Ser	Leu	Leu	Thr	260	265	270	

Val Thr Lys Tyr Tyr Ile Val Pro Tyr Leu Phe Val Asn Phe Trp Leu
 275 280 285

Val Leu Ile Thr Phe Leu Gln His Thr Asp Pro Lys Leu Pro His Tyr
 290 295 300

Arg Glu Gly Ala Trp Asn Phe Gln Arg Gly Ala Leu Cys Thr Val Asp
 305 310 315 320

Arg Ser Phe Gly Lys Phe Leu Asp His Met Phe His Gly Ile Val His
 325 330 335

Thr His Val Ala His His Leu Phe Ser Gln Met Pro Phe Tyr His Ala
 340 345 350

Glu Glu Ala Thr Tyr His Leu Lys Lys Leu Leu Gly Glu Tyr Tyr Val
 355 360 365

Tyr Asp Pro Ser Pro Ile Val Val Ala Val Trp Arg Ser Phe Arg Glu
 370 375 380

Cys Arg Phe Val Glu Asp Gln Gly Asp Val Val Phe Phe Lys Lys
 385 390 395

<210> 21
 <211> 36
 <212> DNA
 <213> synthetic primer

<400> 21
 cuacuacuac uaggatccat ggcacctccc aacact

36

<210> 22
 <211> 41
 <212> DNA
 <213> synthetic primer

<400> 22
 caucaucauc auggtacctc gagttacttc ttgaaaaaga c

41